

Figure 2

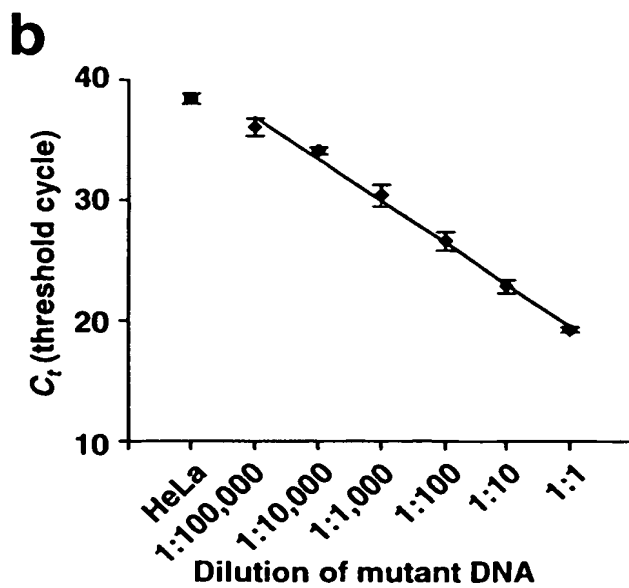
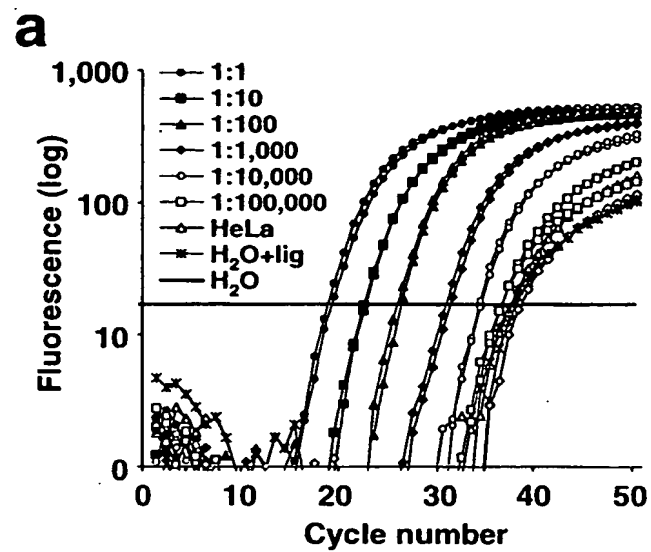


Figure 2 (cont.)

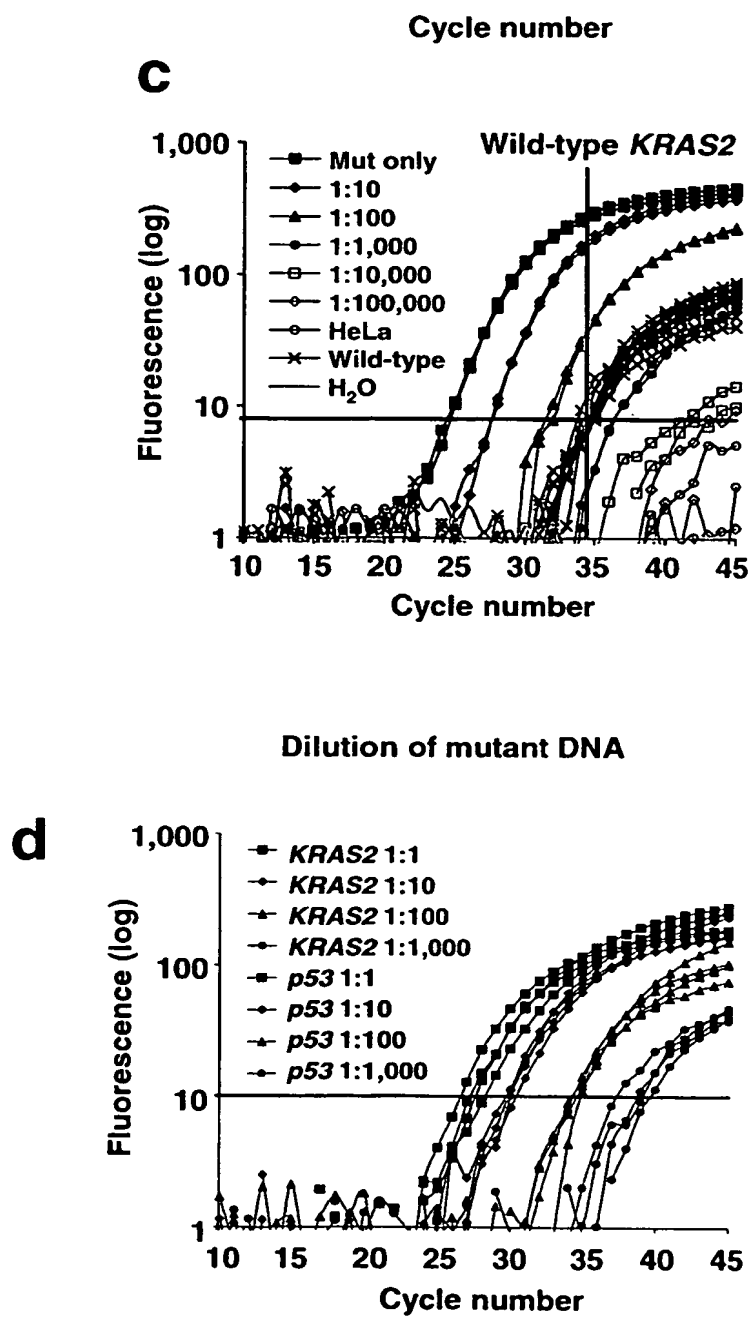


Figure 3

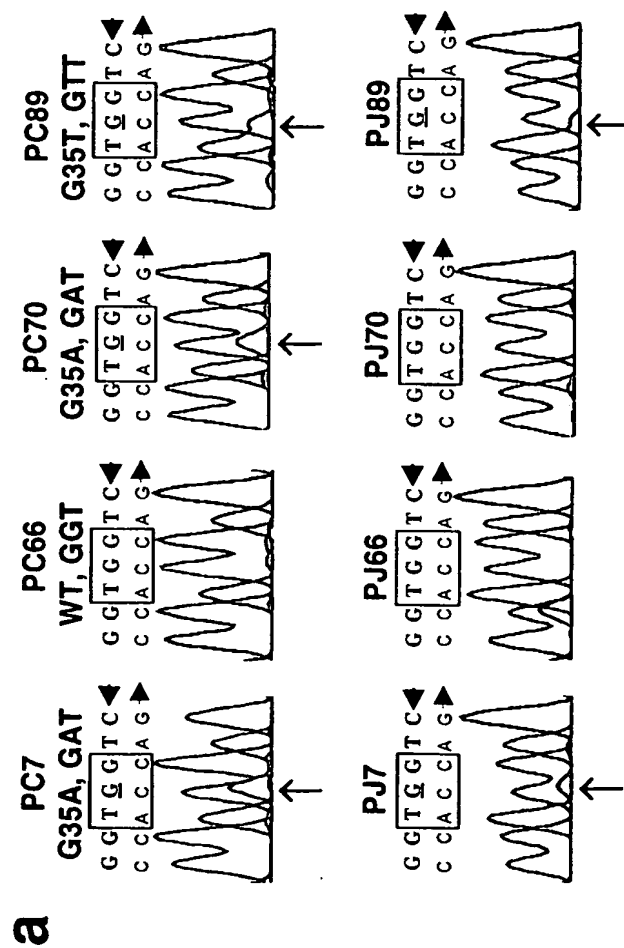


Figure 3 (cont.)

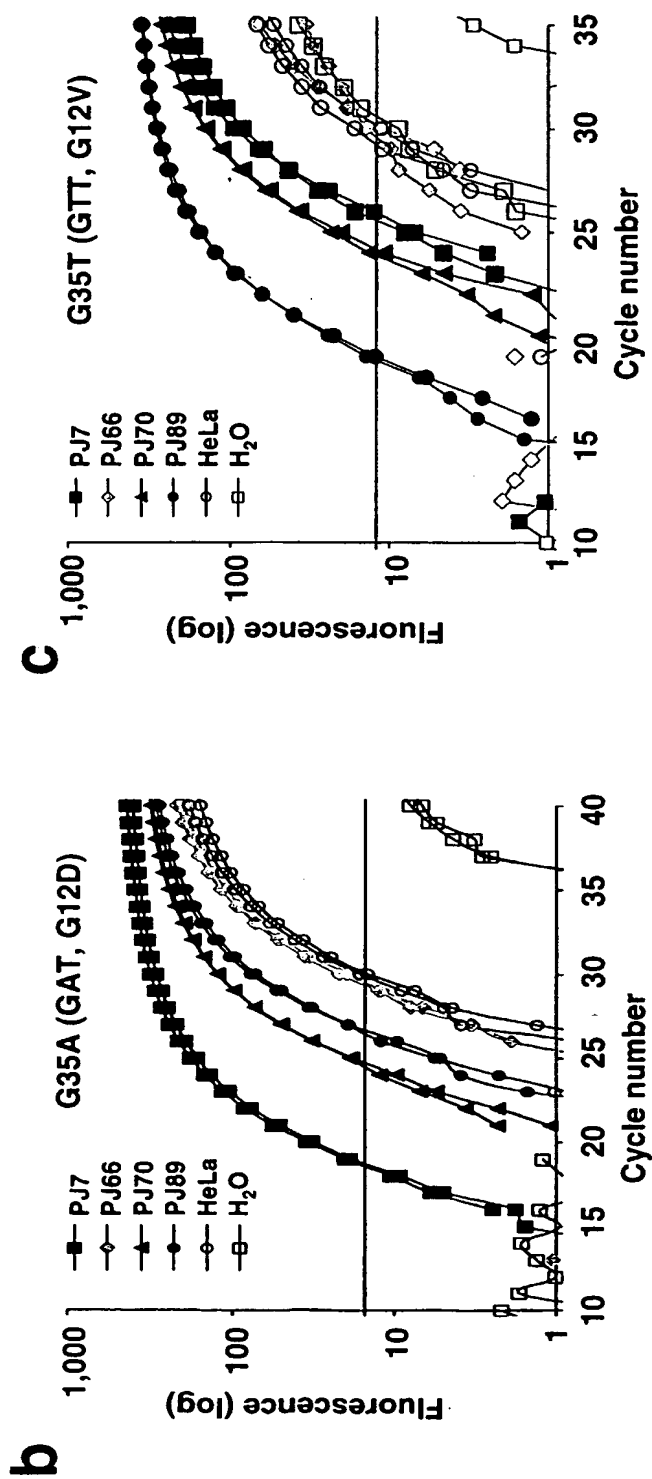


Figure 4

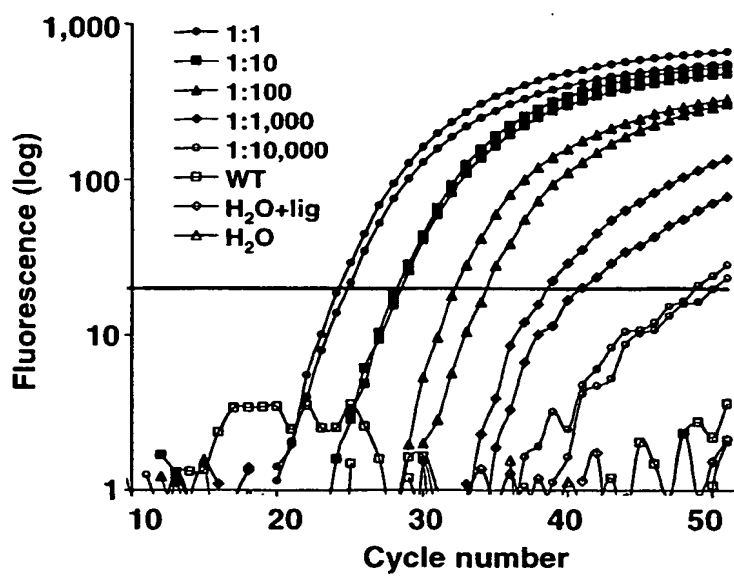
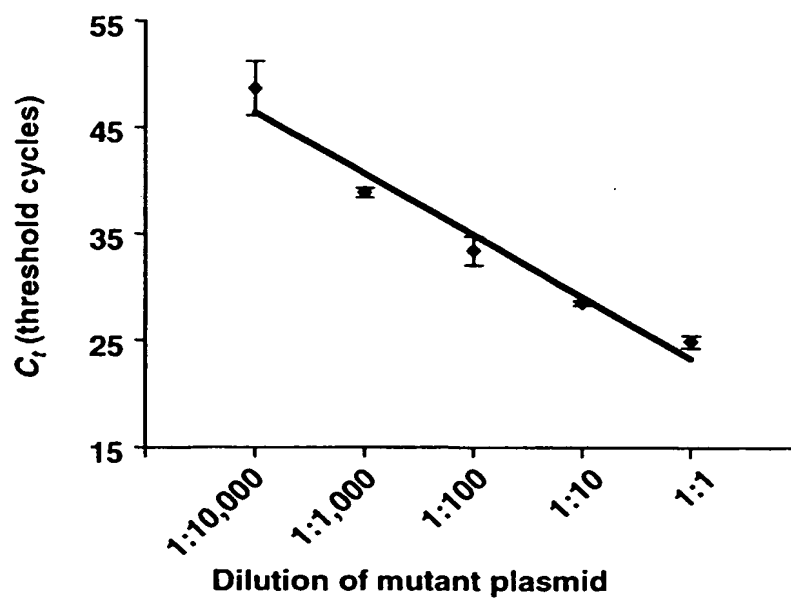
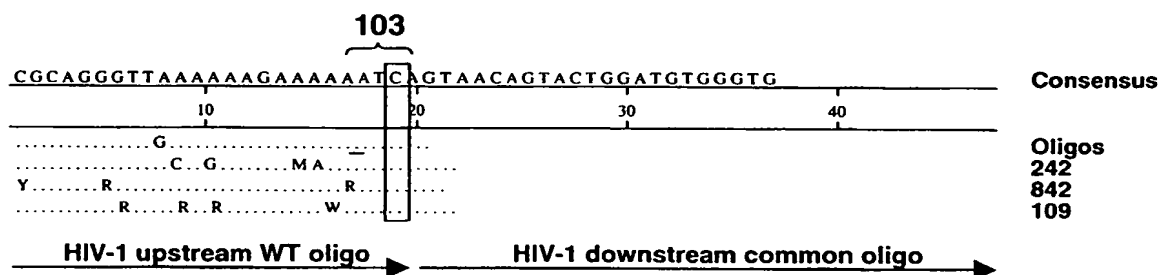
a**b**

Figure 5

a



b

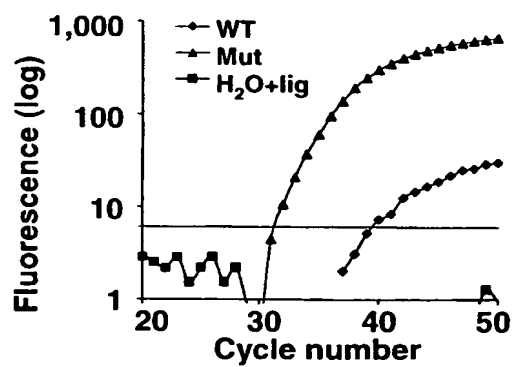
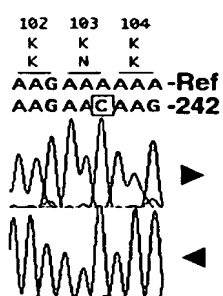


Figure 5 (cont)

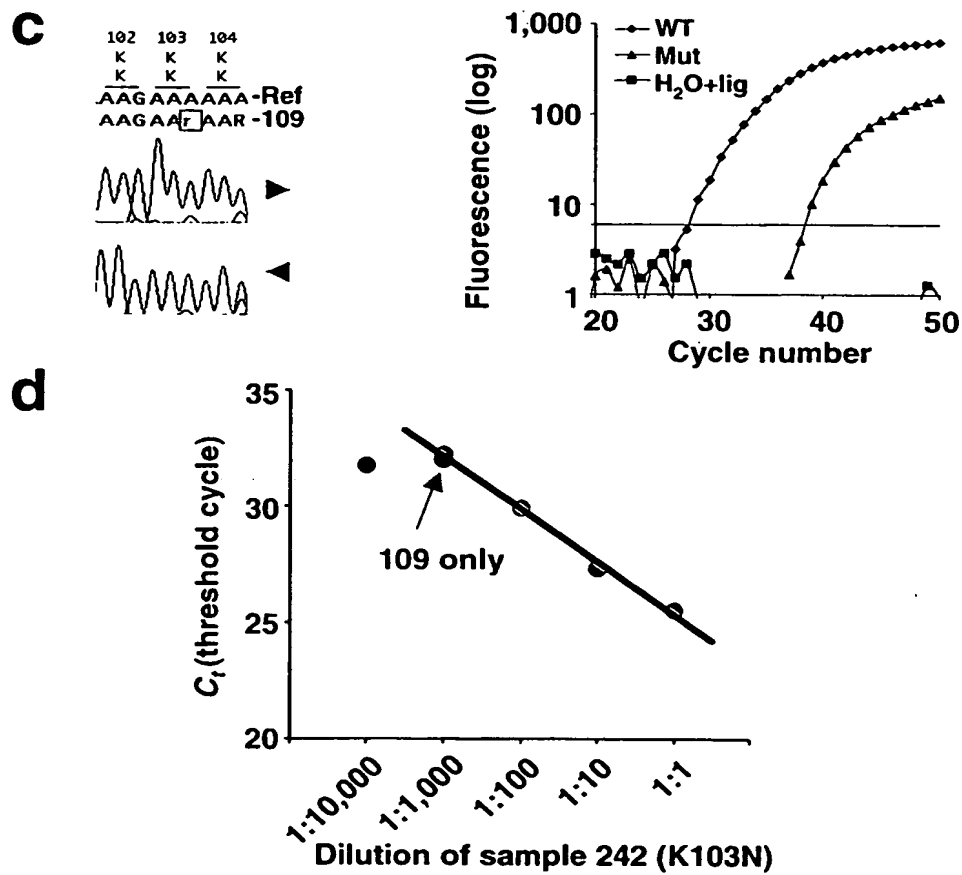


Figure 6

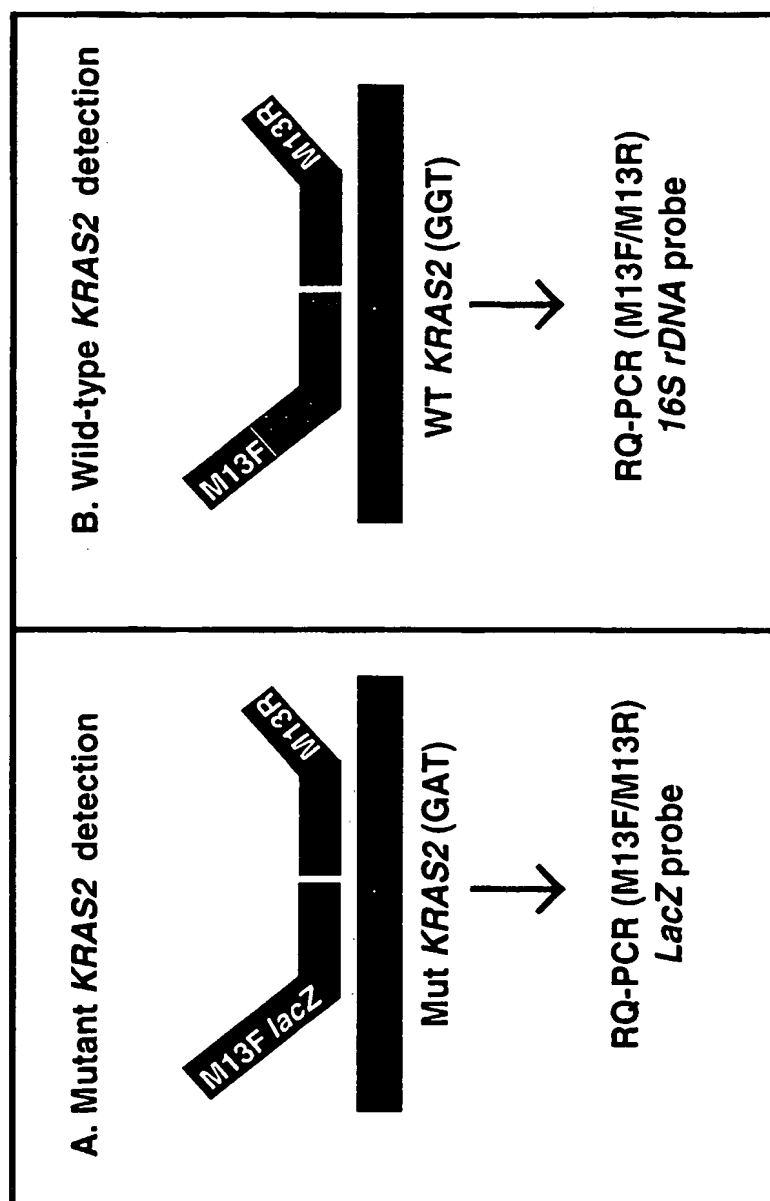


Figure 7A

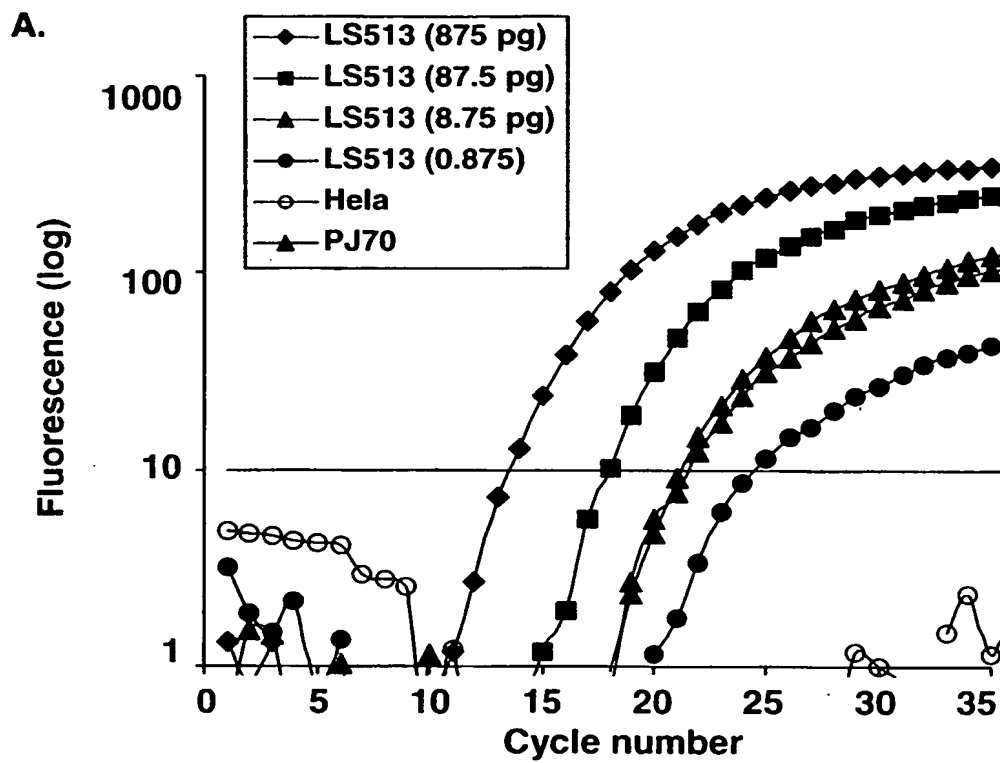
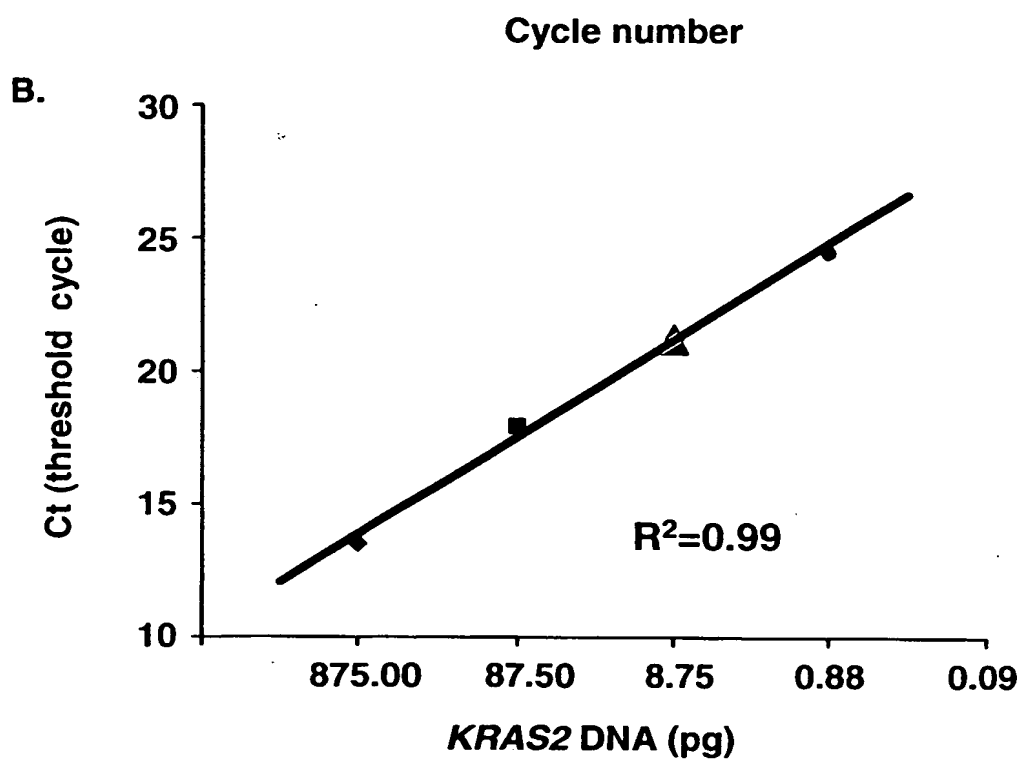


Figure 7B



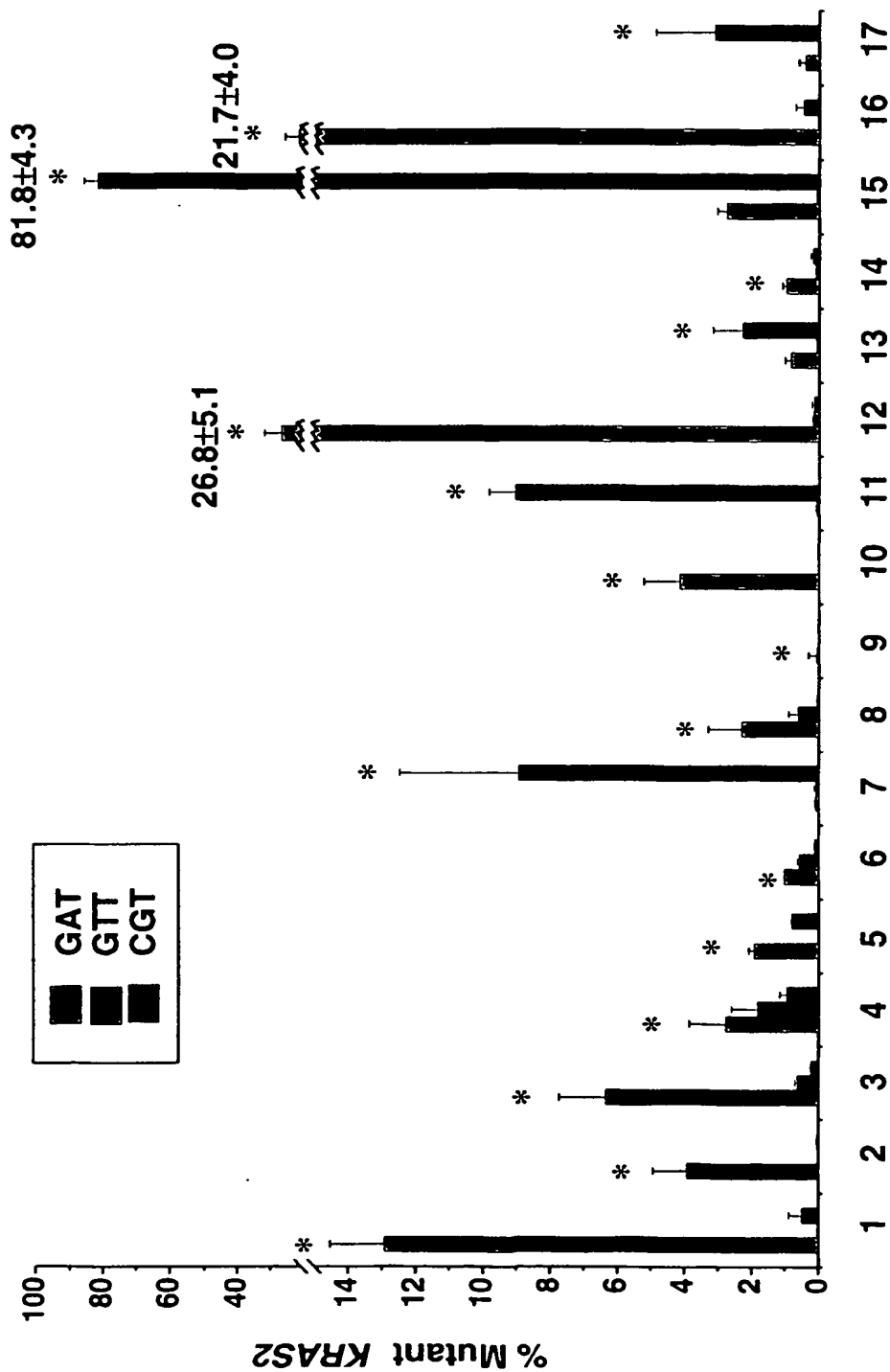


Figure 8

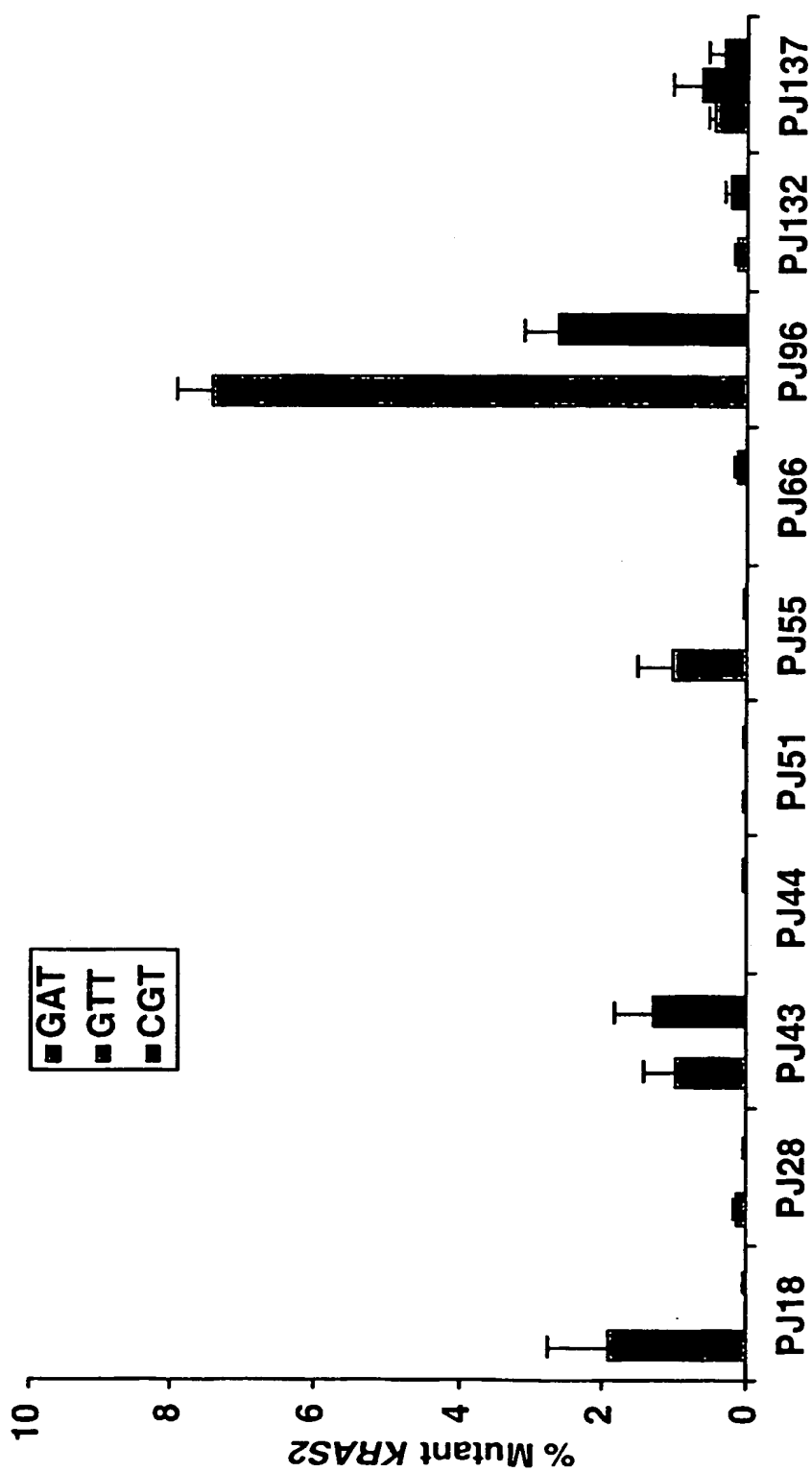


Figure 9

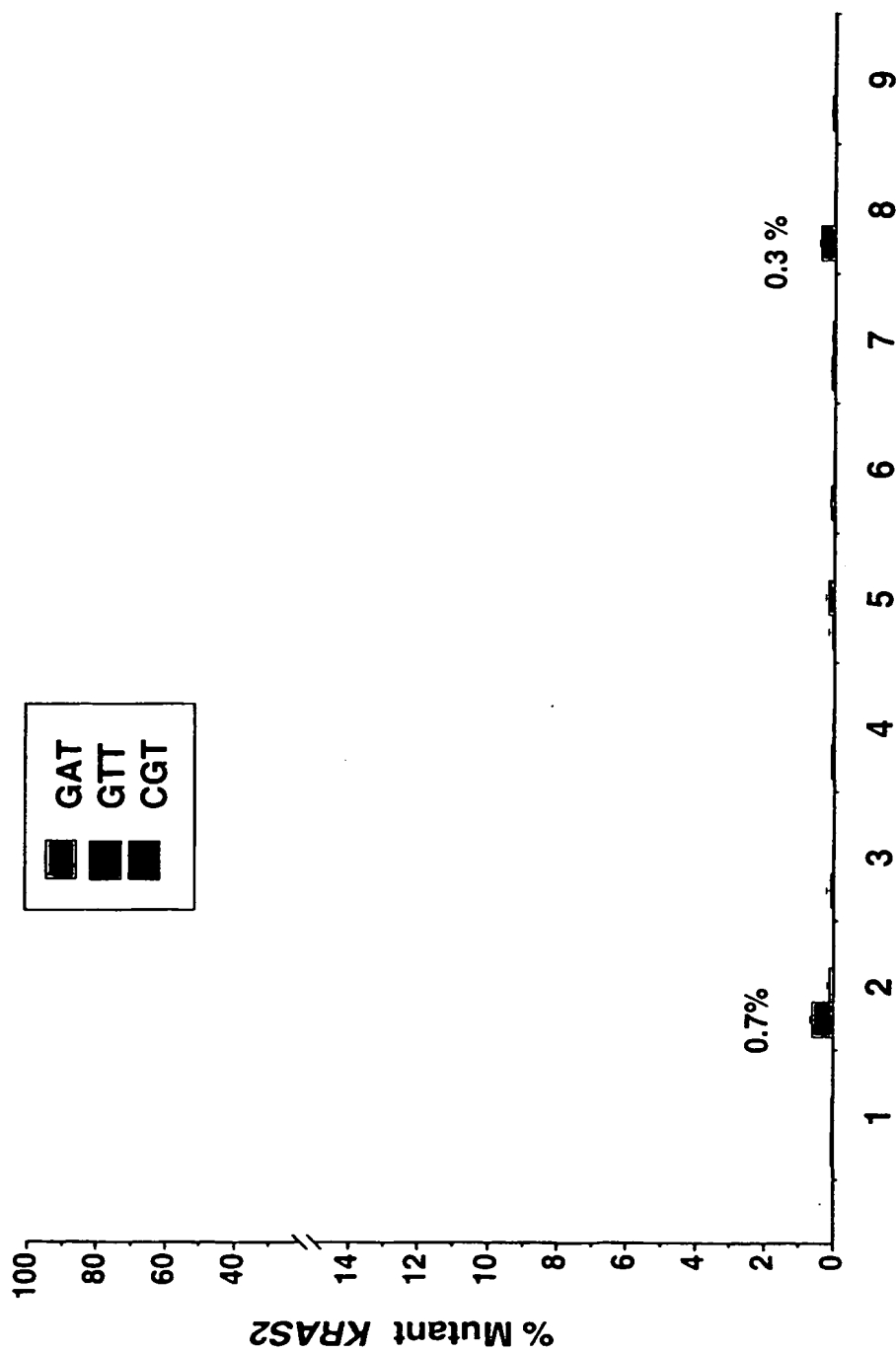


Figure 10

Figure 11

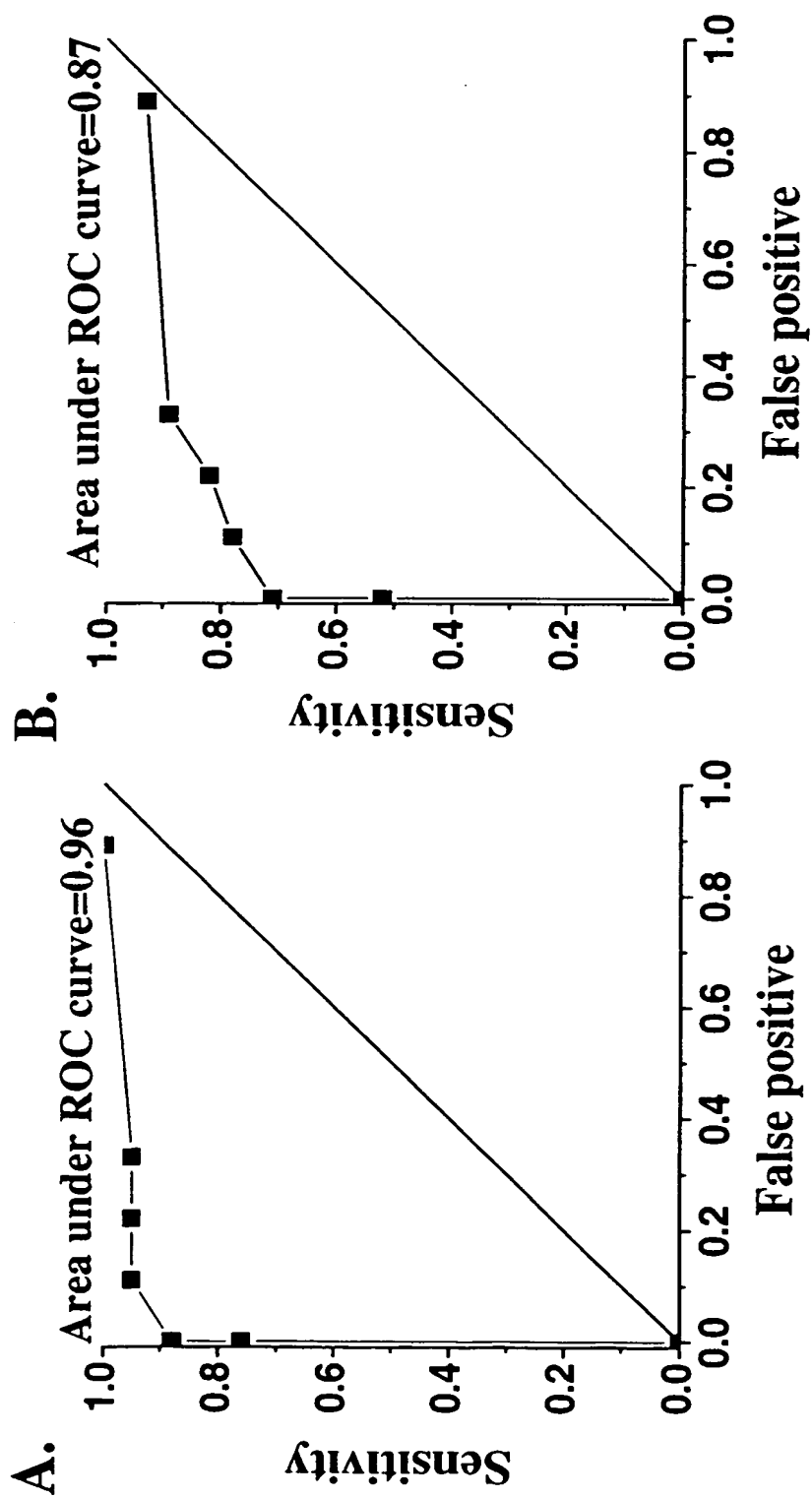


Figure 12

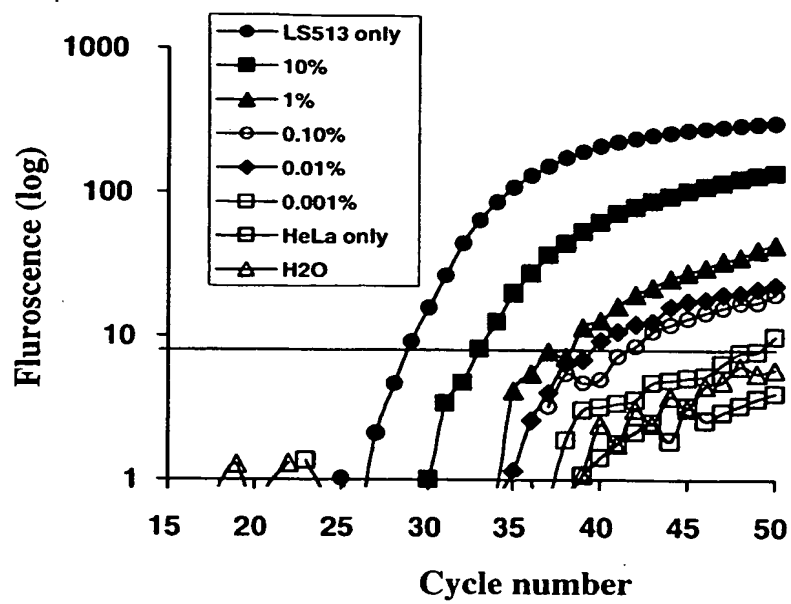


Figure 13

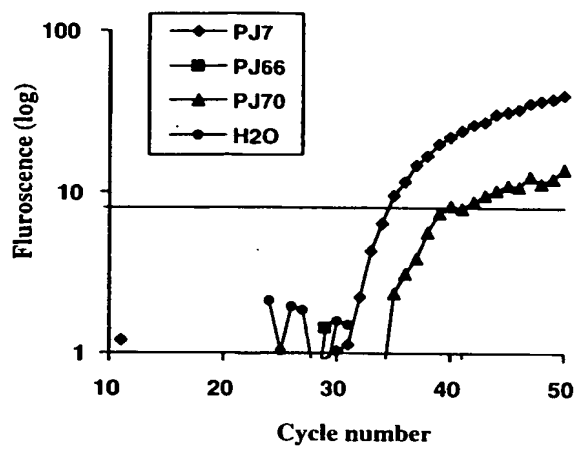


Figure 14

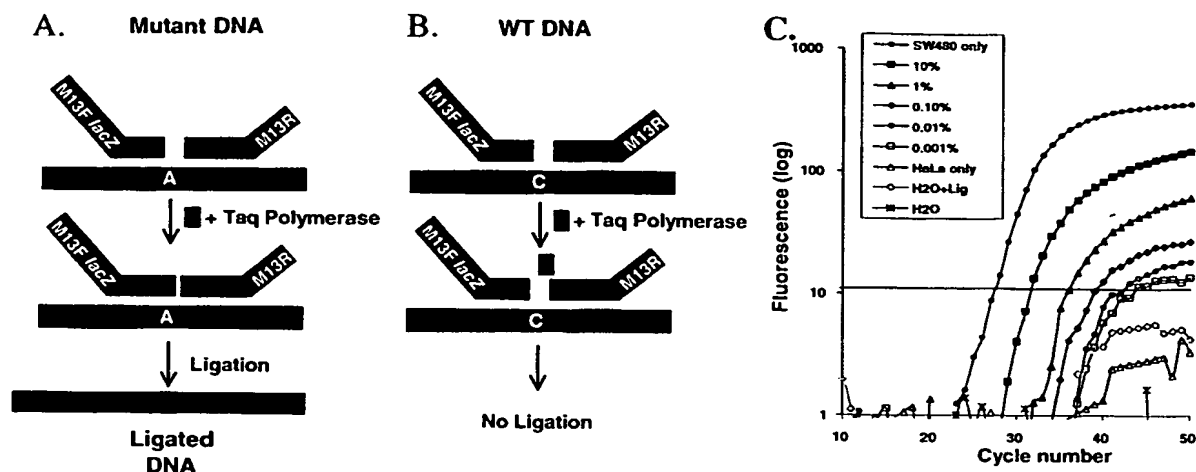


Figure 15

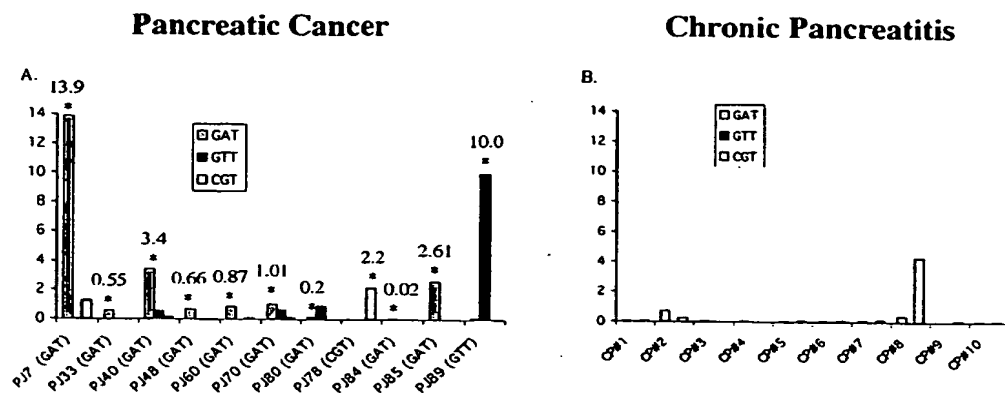


Figure 16

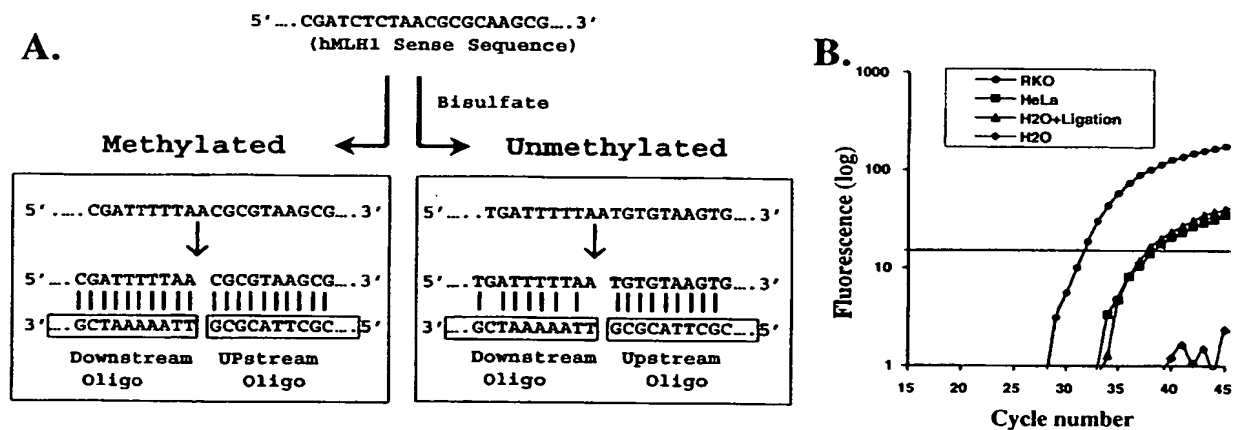


Figure 17

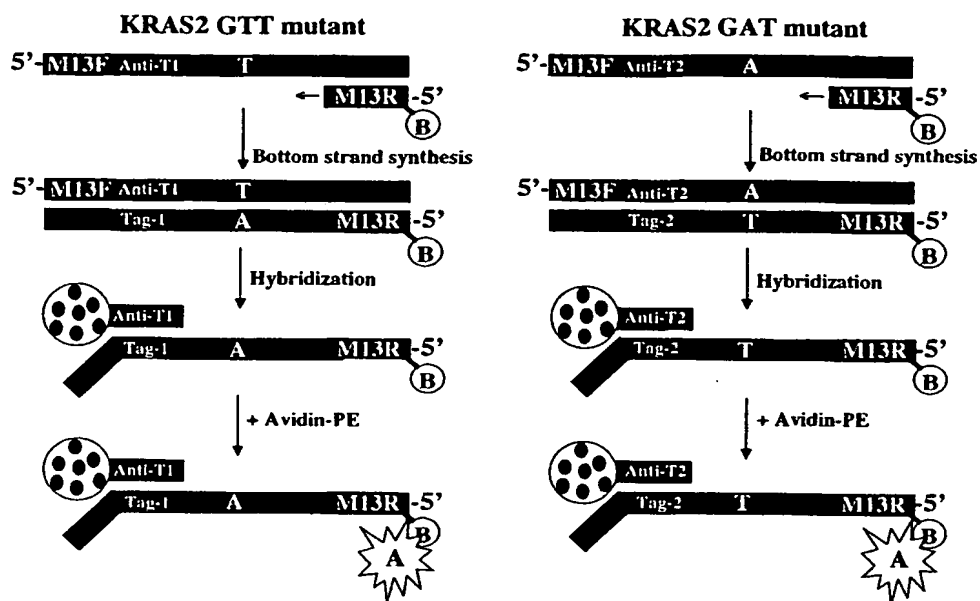


Figure 18

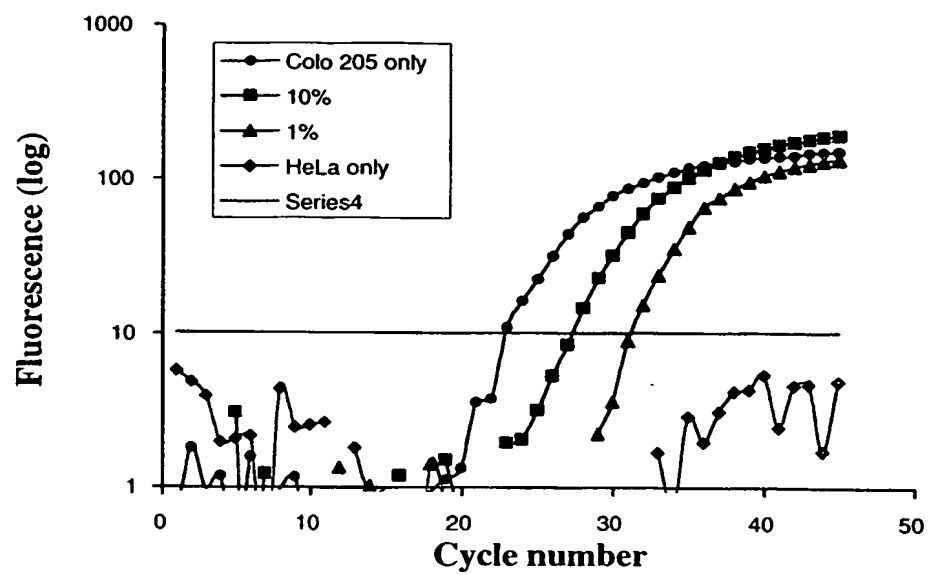
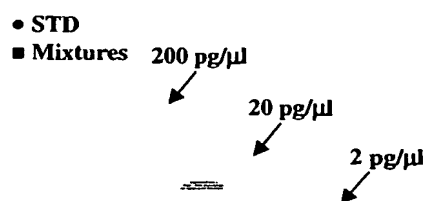


Figure 19

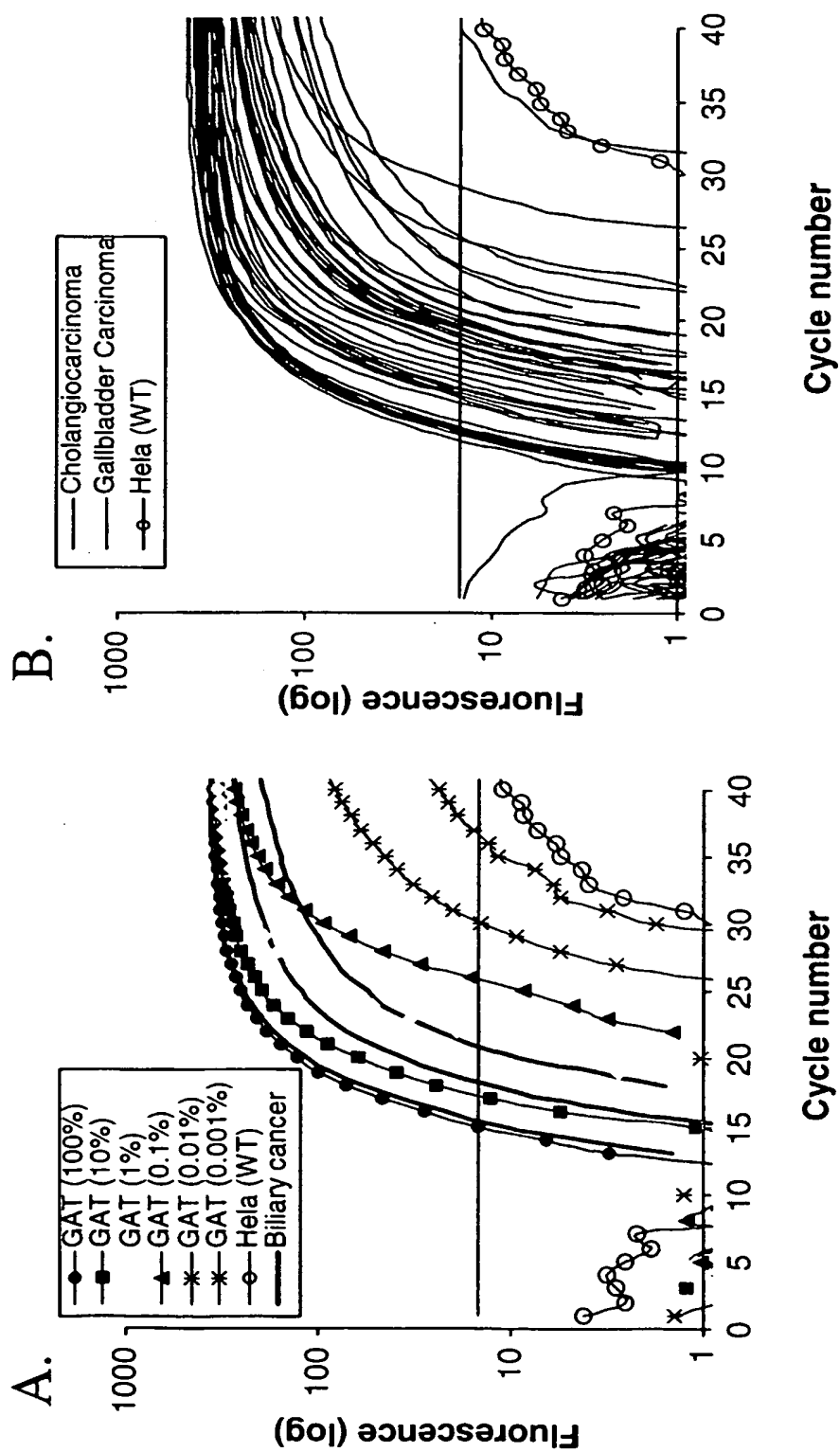
A. Standard curve for quantification



B. % heteroplasmy in the mixtures

Sample #	Heteroplasmy (%)
A	55.3
B	18.7
C	3.3
D	55.7
E	1.1
F	27.5
G	4.02
H	25.6

Figure 20



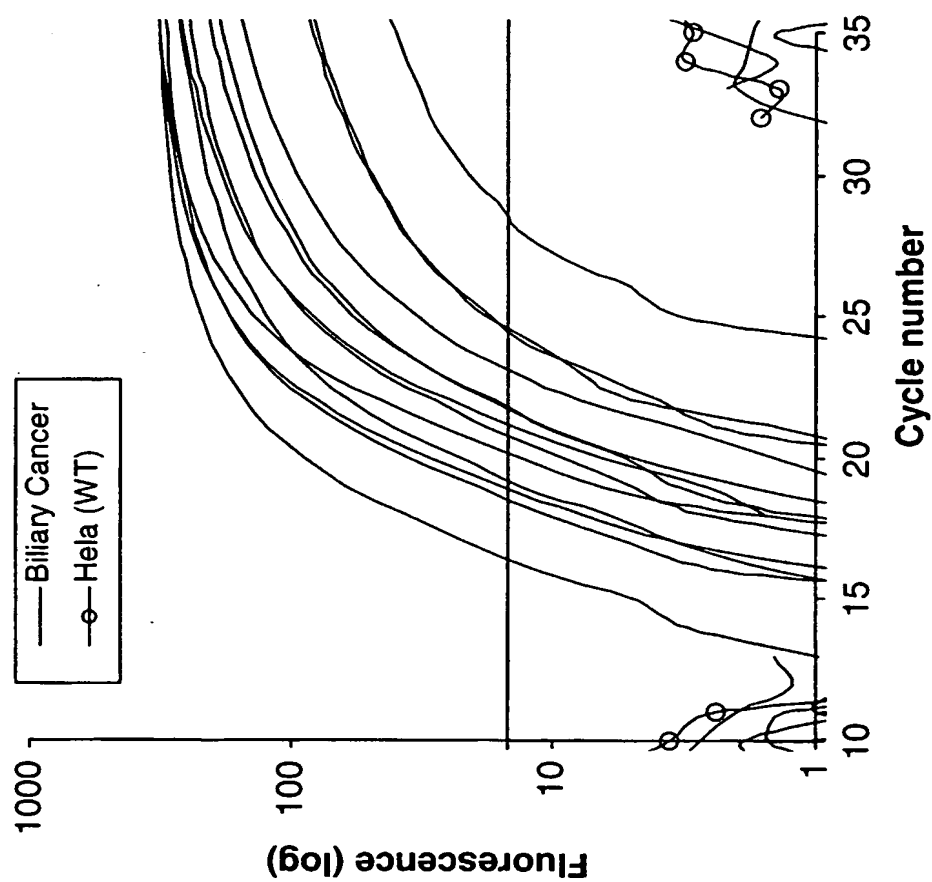
**Figure 21**

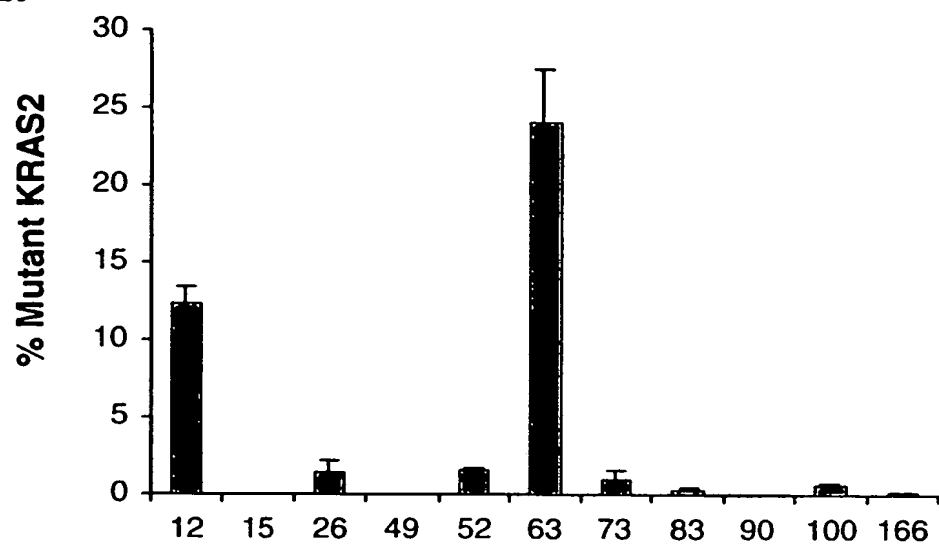
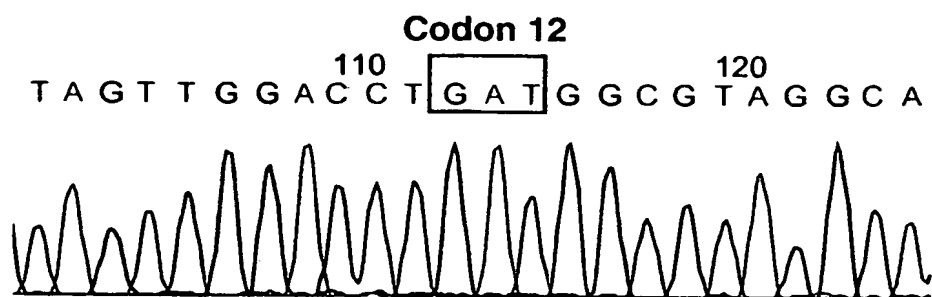
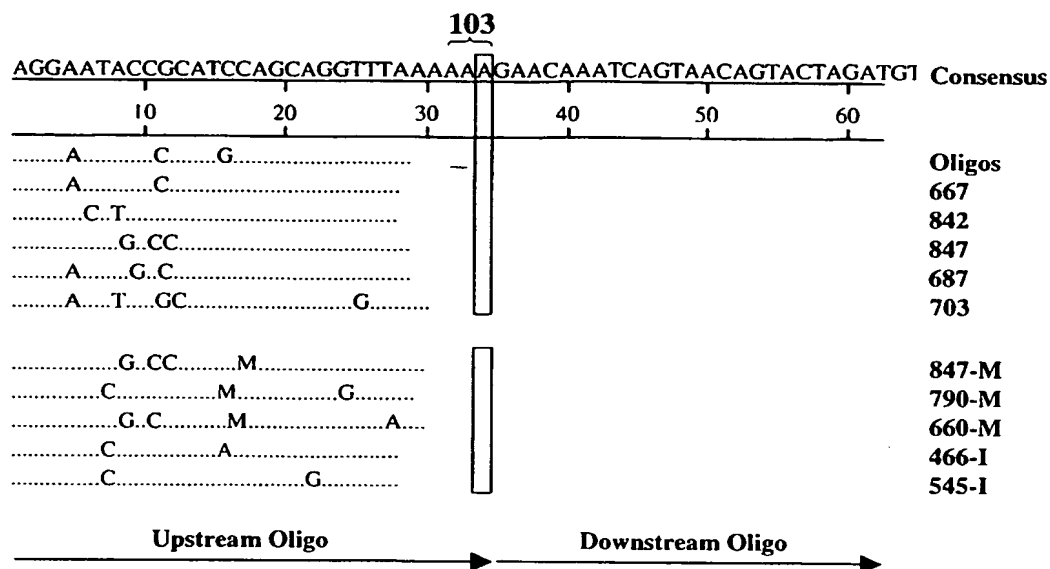
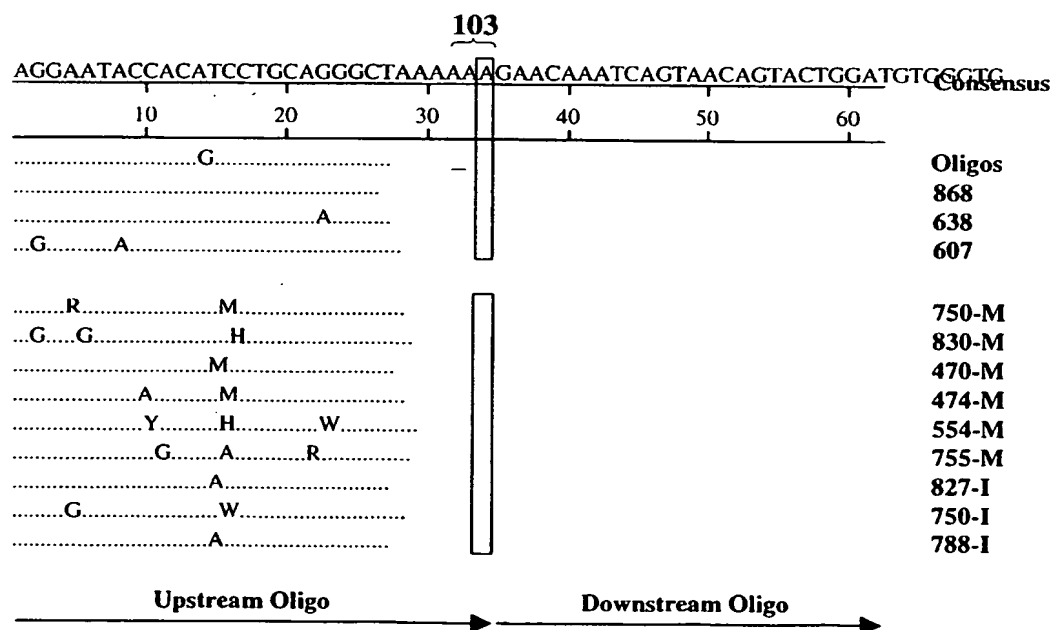
Figure 22**A.****B.**

Figure 23

A**B**

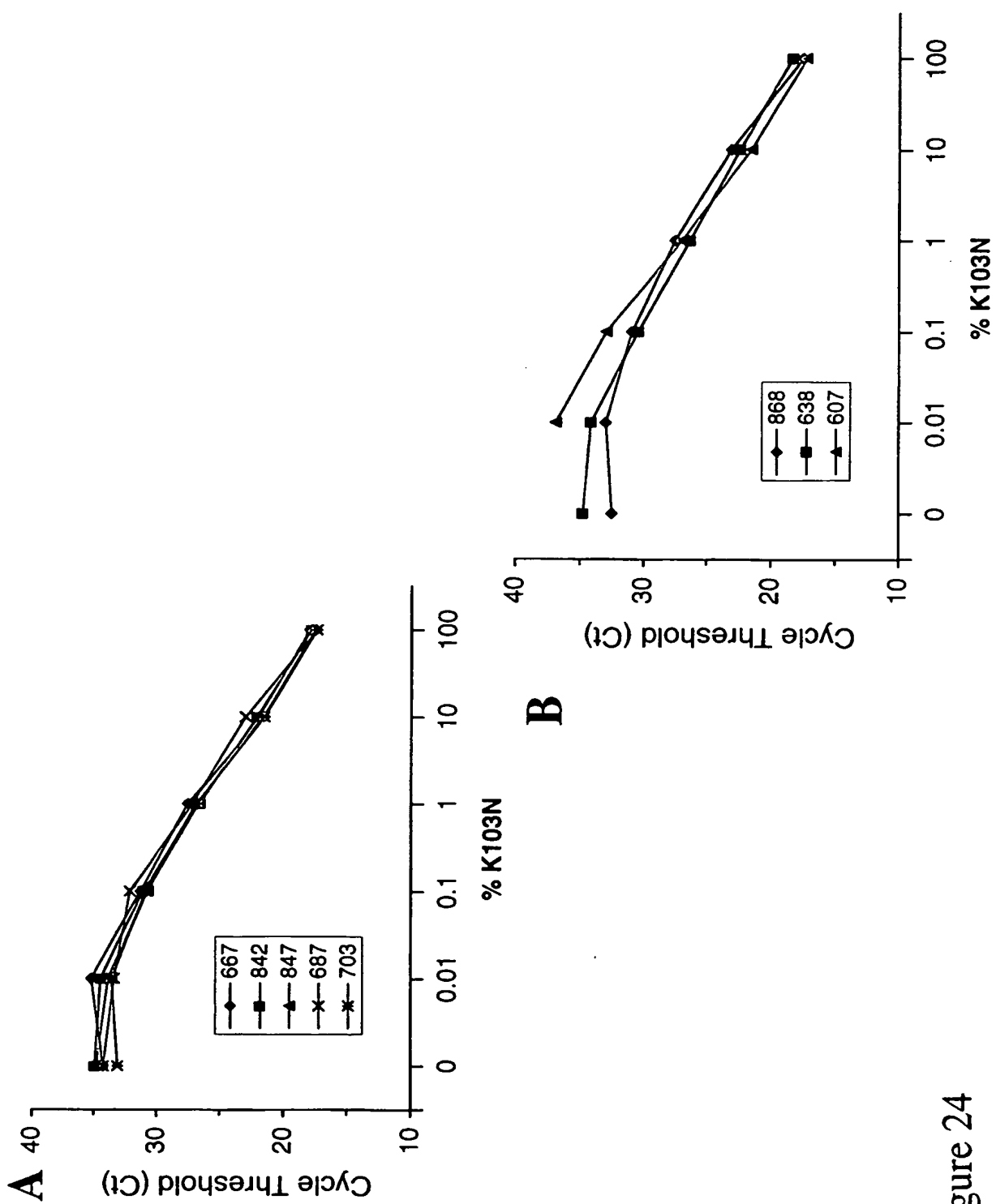


Figure 24

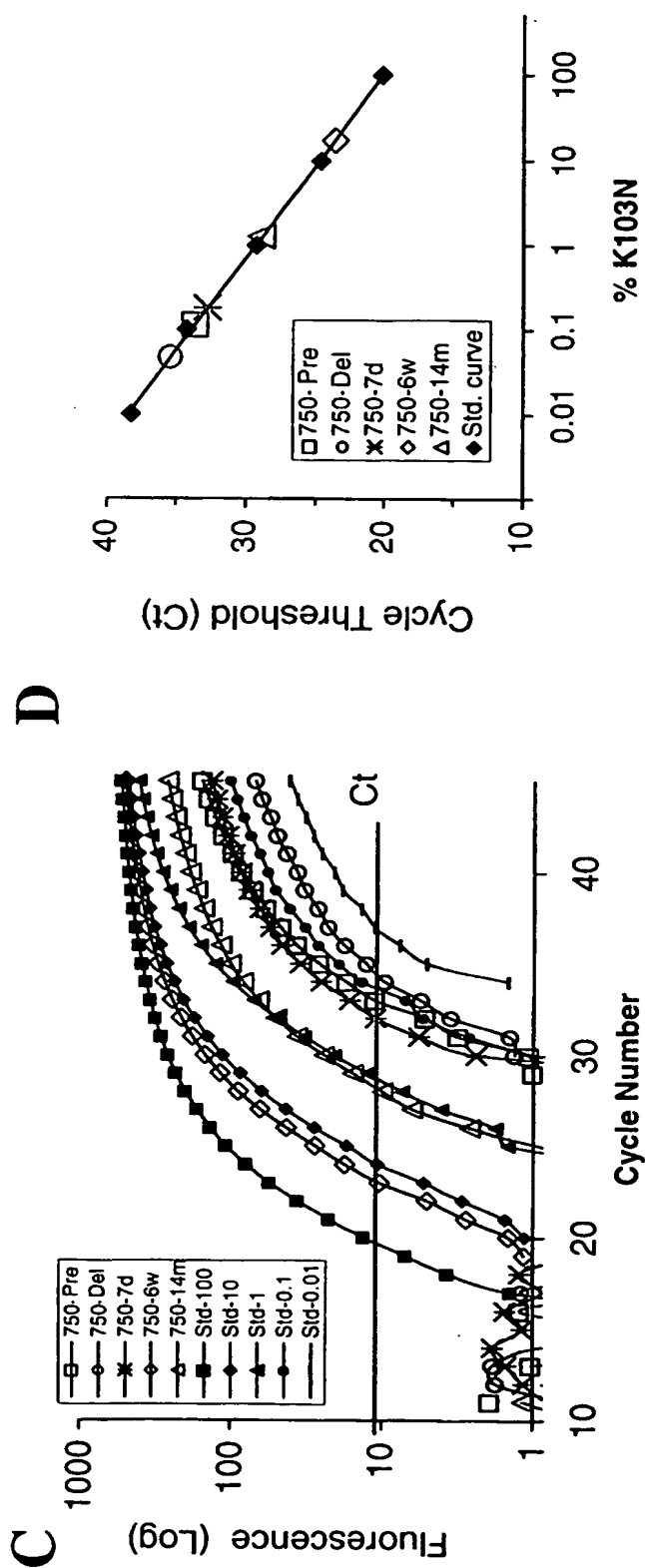
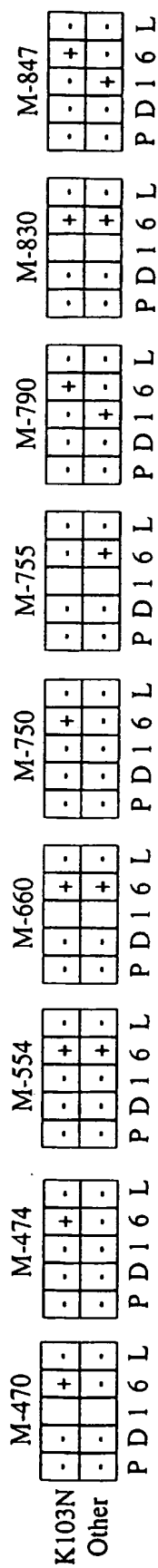
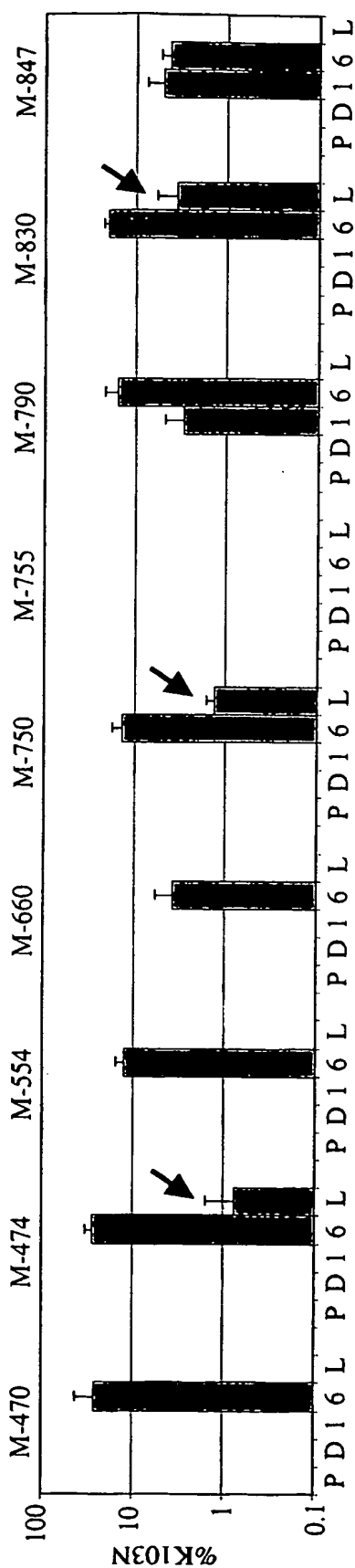


Figure 24 (cont.)

A ViroSeq



B. LigAmp



C TyHRT

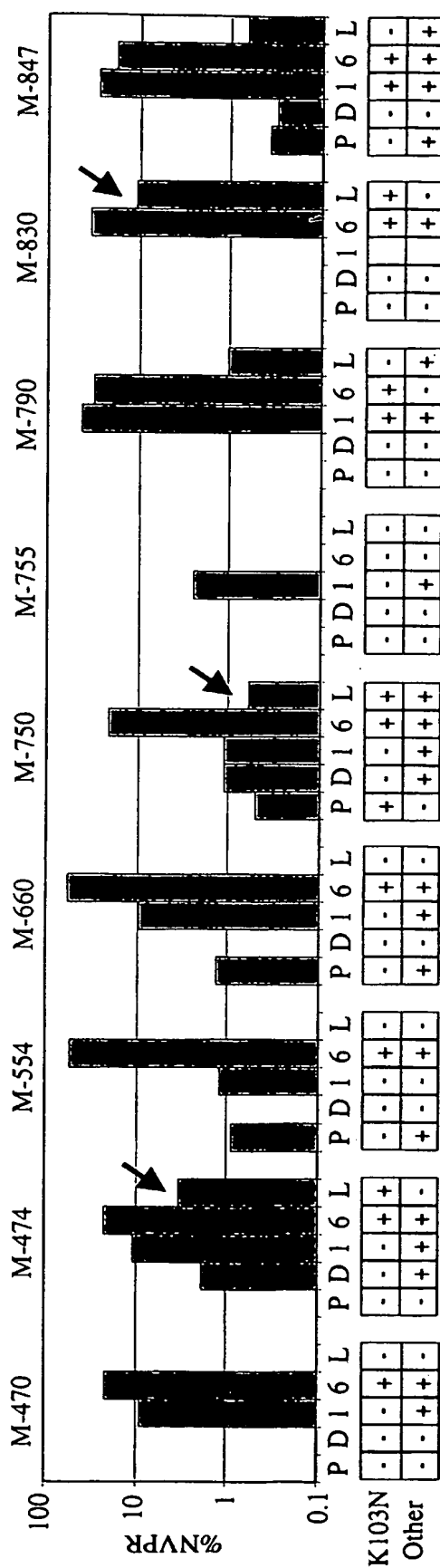


Figure 26

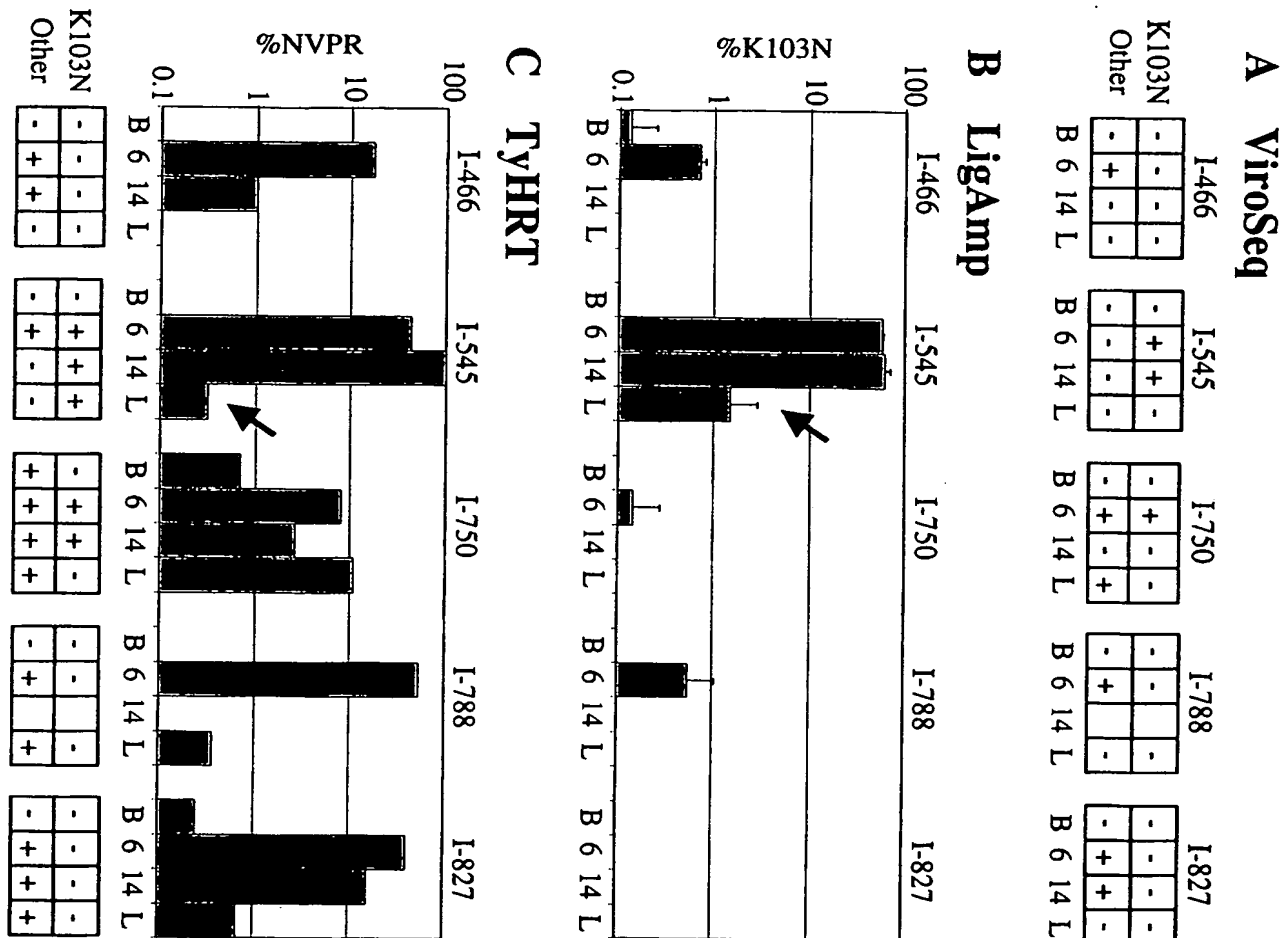
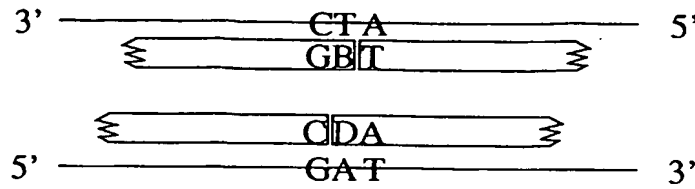
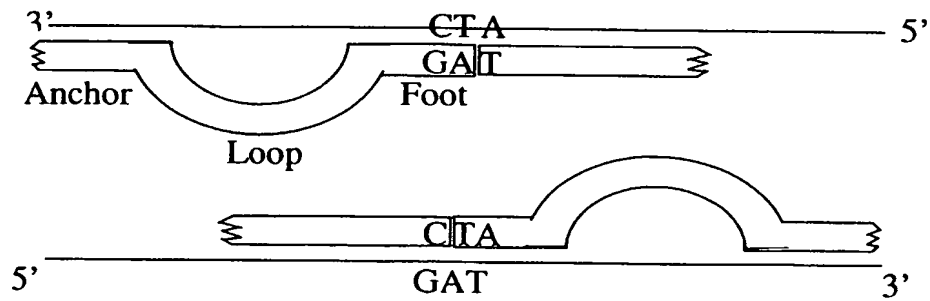


Figure 27

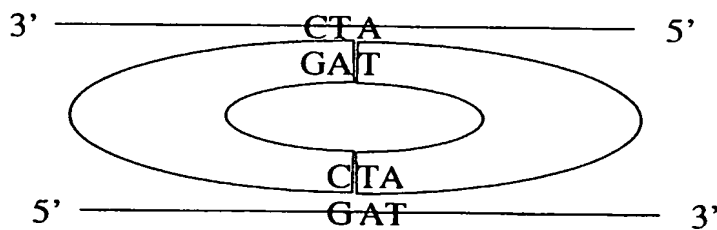
A.



B.



C.



D.

